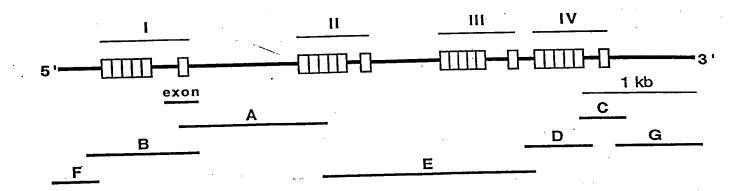


1



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Fig. 2



NAIDM	7
538ge para	
GSFPPELA	GSFPPELASTPLEDIDPFYSNVLTFVVISKGKDIFRFSASKAMWLLDPFNPIRRVAIYILVHPLFSLFIITTILTNCILMIMPTTPTVESTEVIFTGIYT 179
FESAVKVM	IS3 IS4 IS5 FESAVKVMARGFILCPFTYLRDAWNWLDFVVIALAYVTMGIDLGNLAALRTFRVLRALKTVAIVPGLKTIVGAVIESVKNLRDVIILTMFSLSVFALMGL 279 279
OIXMGVLT	P
EDLYQHVL H	1 S 6 EDLYQHVLQAAGPWHMLFFIVIIFLGSFYLVNLILAIVAMSYDELQKKAEEEEAAEEEAIREAEEAAAAKAAKLEERANVAAQAAQDAADAAAALHPEM 479HQVAQ
AKSPTYSC	* AKSPTYSCISYELFVGGEKGNDDNNKEKMSIRSVEVESESVSVIQRQPAPTTAP.ATKVRKVSTTSLSLPGSPFNLRRGSRSSHKYTIRNGRGRFGIPGS 578
DRKPLVLQ Q S	* DRKPLVLQTYQDAQQHLPYADDSNAVTPMSEENGAIIVPAYYCNLGSRHSSYTSHQSRISYTSHGDLLGGMAAMGASTWTKESKLRSRNTRNQSIGAATN 678QQ

## Fig. 3 (contil)

GGSSTAGGGYPDANHK. EQRDYEMGQDYTDEAGKIKHHDNPFIEPVQTQTVVDMKDVMVIANDIIEQAAGRHSRASERGEDDDEDGPTFKDIALEYILKGI 777  -GSSTAGGGYP-A EQM-QDY	WDCMYVGDVSC	SEQPSEHCONELELGHDEINGEGLIKKGMKGETQLEVAIGDGMEFTIHGDMKNNKPKKSKFMNNTTMIGNSINHQDNRLEHELNHRGLSIQUEDDTASIN 1177
GGSSTAGGGYPDANHKGSSTAGGGYP-ATTCLTI EIFCVWDCCWWLKFQ! EI DV	11P VFRVLCGEWIESMWDCM	ISDQPSEHGDNELELGI  SYGSHKNRPFKDESHK  LRLKTFQLIENKYFETY  LRLKTFQLIENKYFETY  IIIS4  ND-AV-RS GG-QA-KT

E.

Fig. 4